

FIG. 1A

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1 GTTTTCAAAA TCGGTGGAGG TGCAATGAAAA AGTTATTGGG CATTTTTGAA AAATAAAAAA
61 ATATCAATAA GTTGGAGTC TTACCGAATT TTTATACCTTA TTTGTTTAGA ATGAACTTA
-35 -10
121 TAACATAGTT GGATAGAGTT TTGATTTAA TACATTAAT GTGAACCTTG CTACAACAAG
181 ATGTGCATCA GAAGGAGTGG TTTAATAATC --sarR--TAA
Ribosomal binding site [SEQ. ID. No. 1]

FIG. 1B

	10	20	30	40	50			
SarR	M--SKINDIDLVNATFQVKFFRDTKKFKNLNYEEIYILNHILRSESNEISSKELAKCS							
SarA	MAITKINDCFELLSMVTYADKLKSLIKKEFSISFEFAVLTYISENKEKEYYLKDIINHL							
		60	70	80	90	100	110	
SarR	EFKPYYLTKAQQLKDLKLLSQKRILQDERTVIVYVT-----DTLKANIHKLISELEE							
SarA	NYKQPQVVKAVKILSQEDYFDKKRNEHDERTVLILVNAQQRKKIESLLSRVNKRITEANN							

SarR YIKN

SarA EIEL

SarR = SEQ ID No. 2

SarA = SEQ ID No. 3

FIG. 1C

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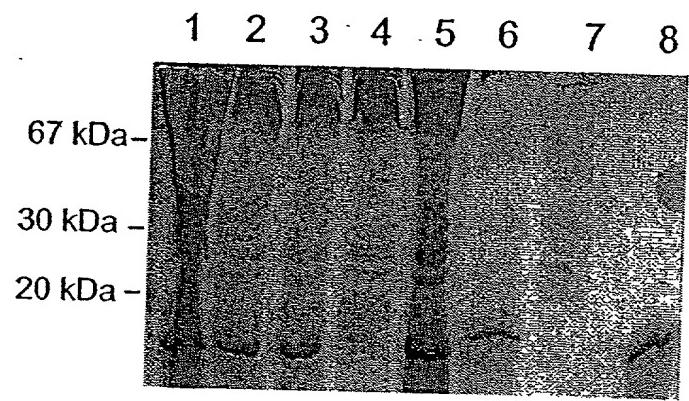


FIG. 2

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FIG. 3A PCR

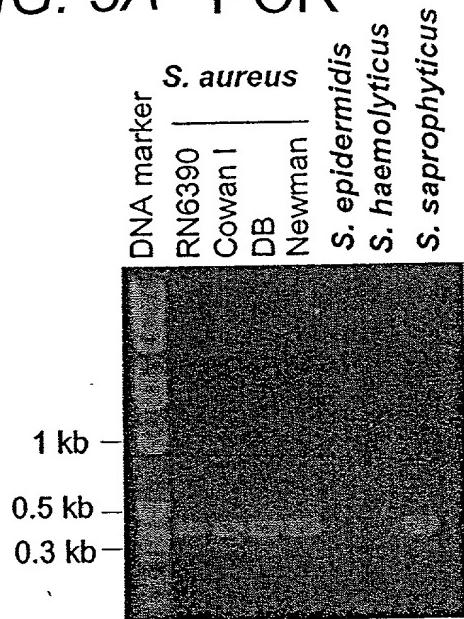


FIG. 3B Southern

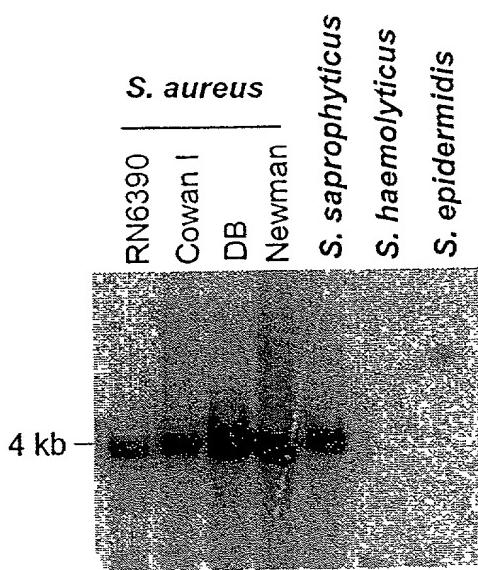


FIG. 3C Northern

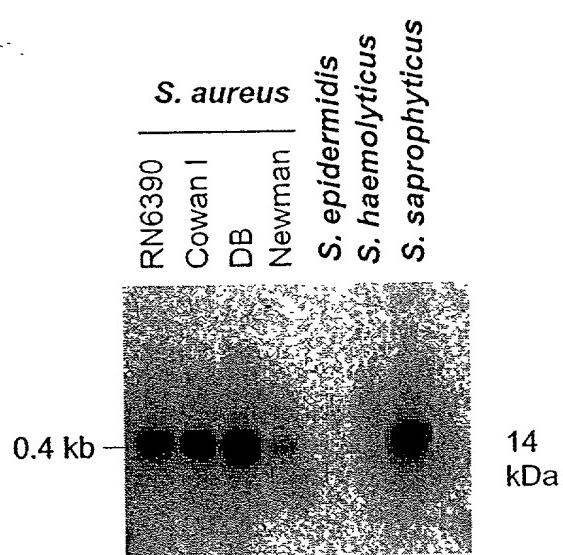
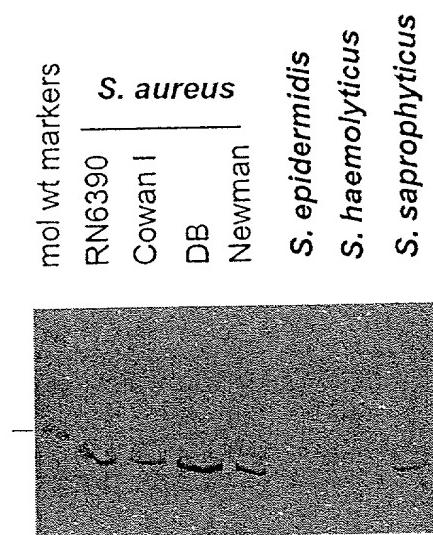


FIG. 3D Immunoblot



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FIG. 4A

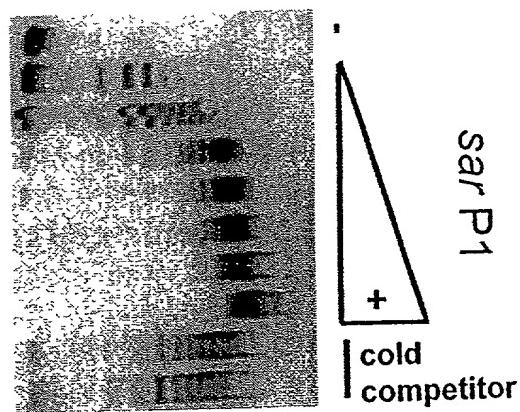


FIG. 4B

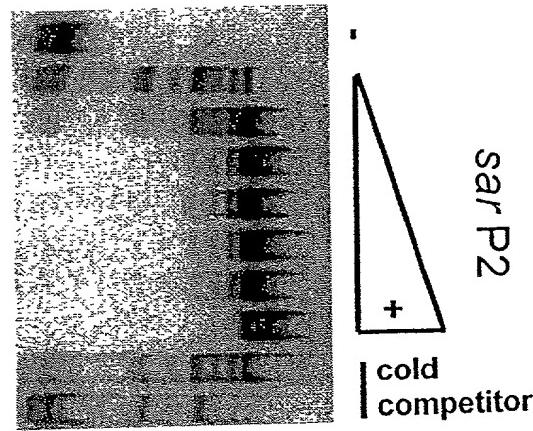
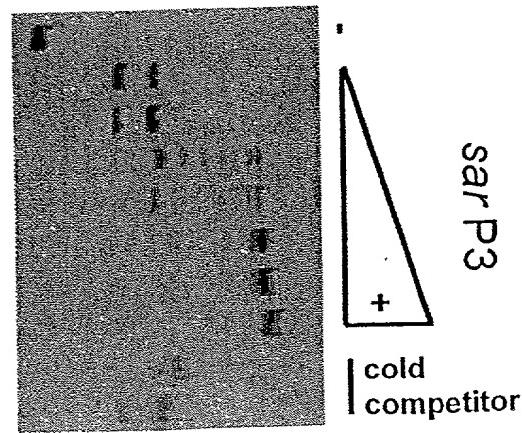
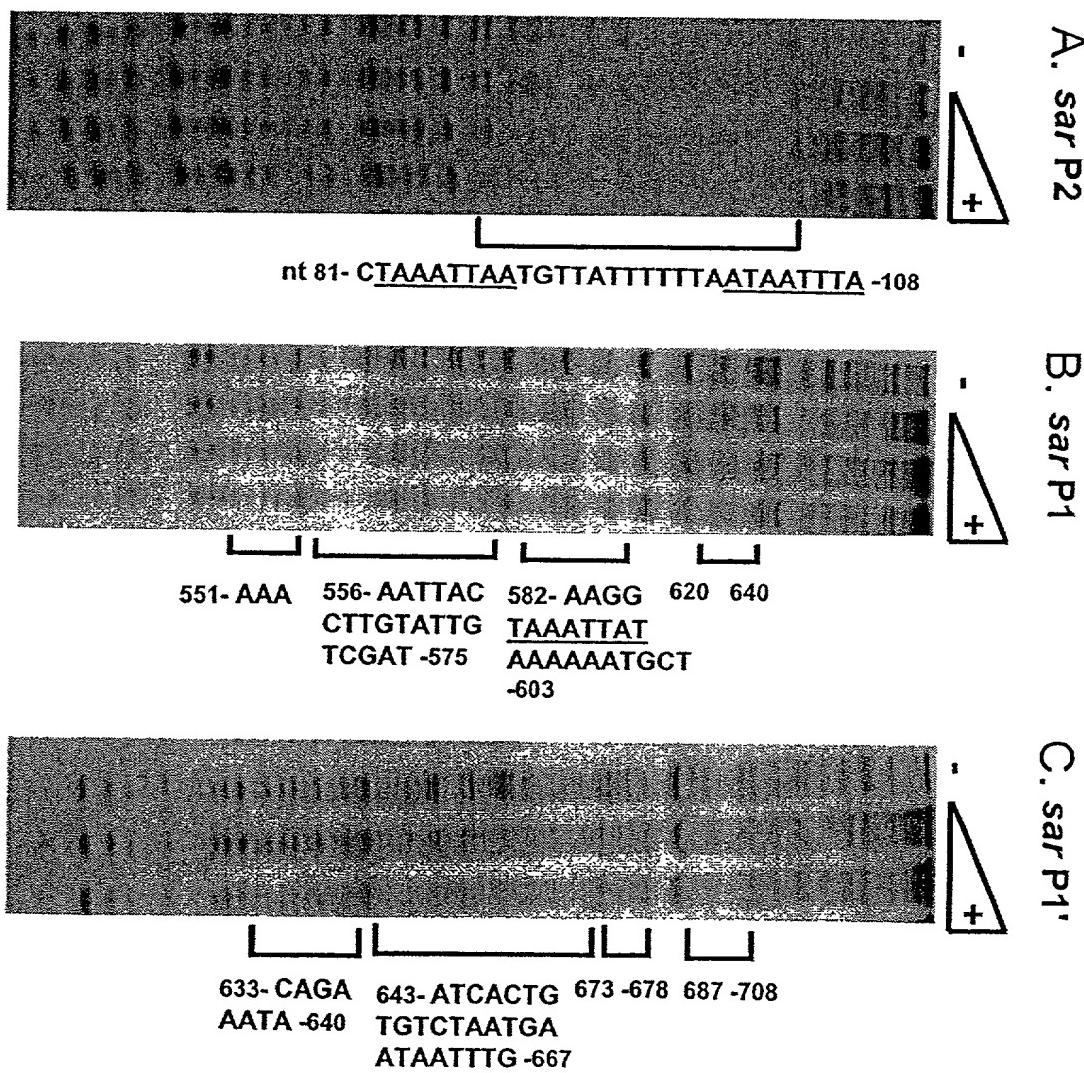


FIG. 4C



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FIG. 5



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FIG. 6A

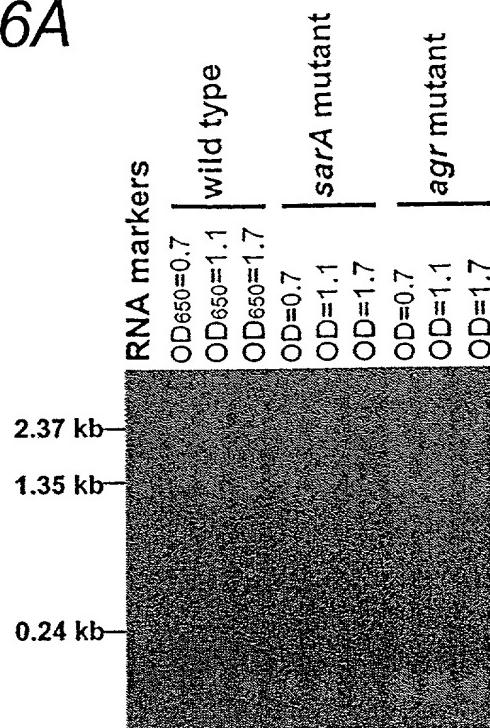


FIG. 6B

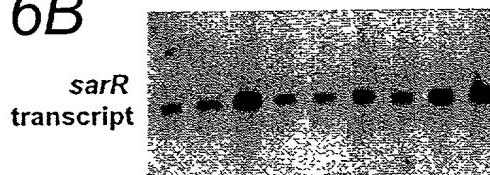


FIG. 6C

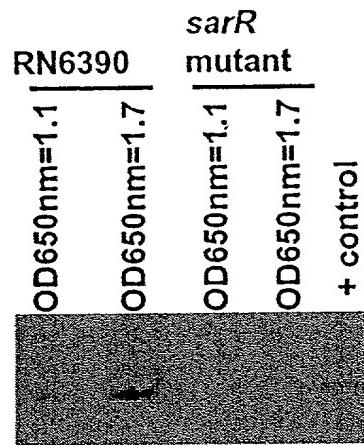


FIG. 7A

sar P1

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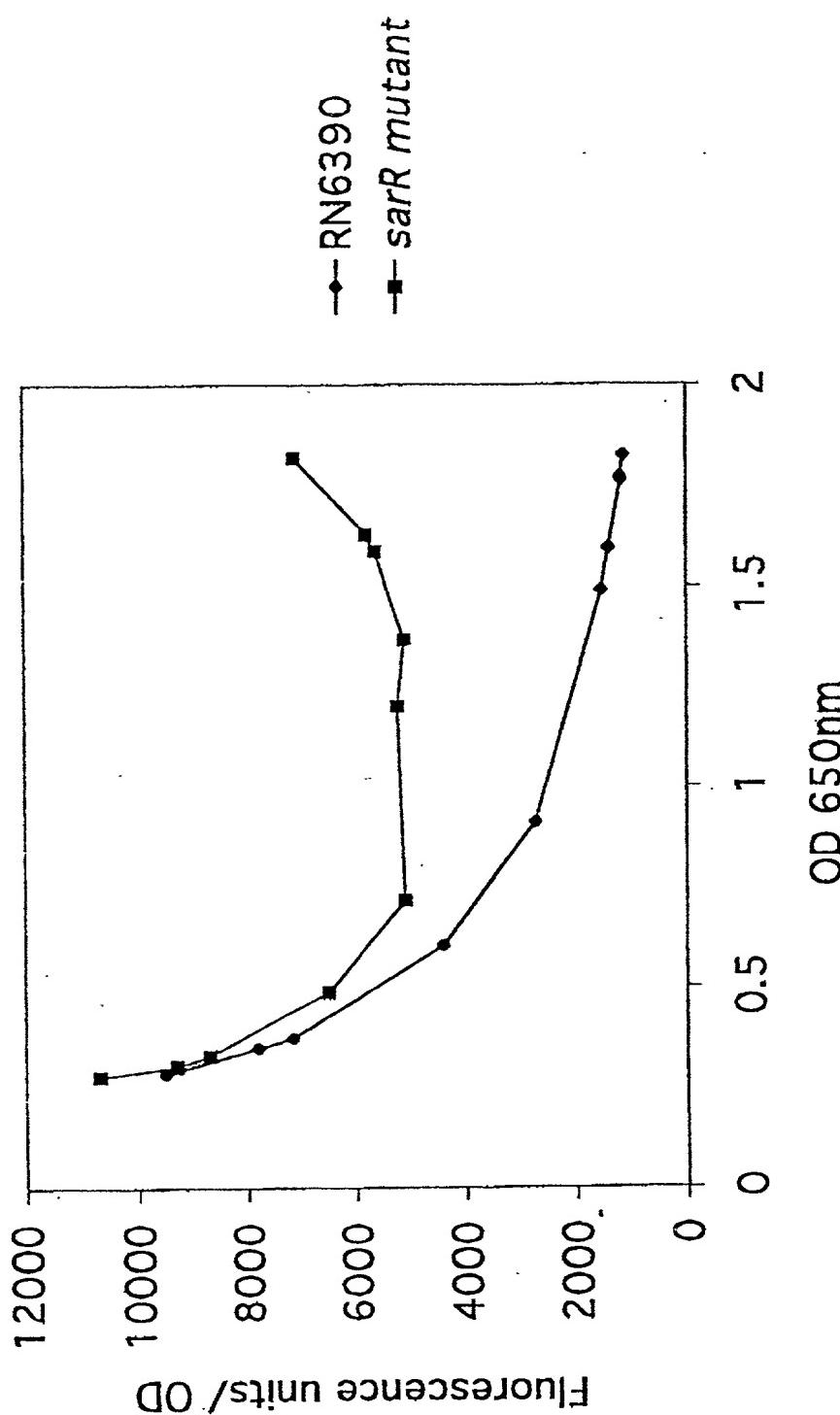
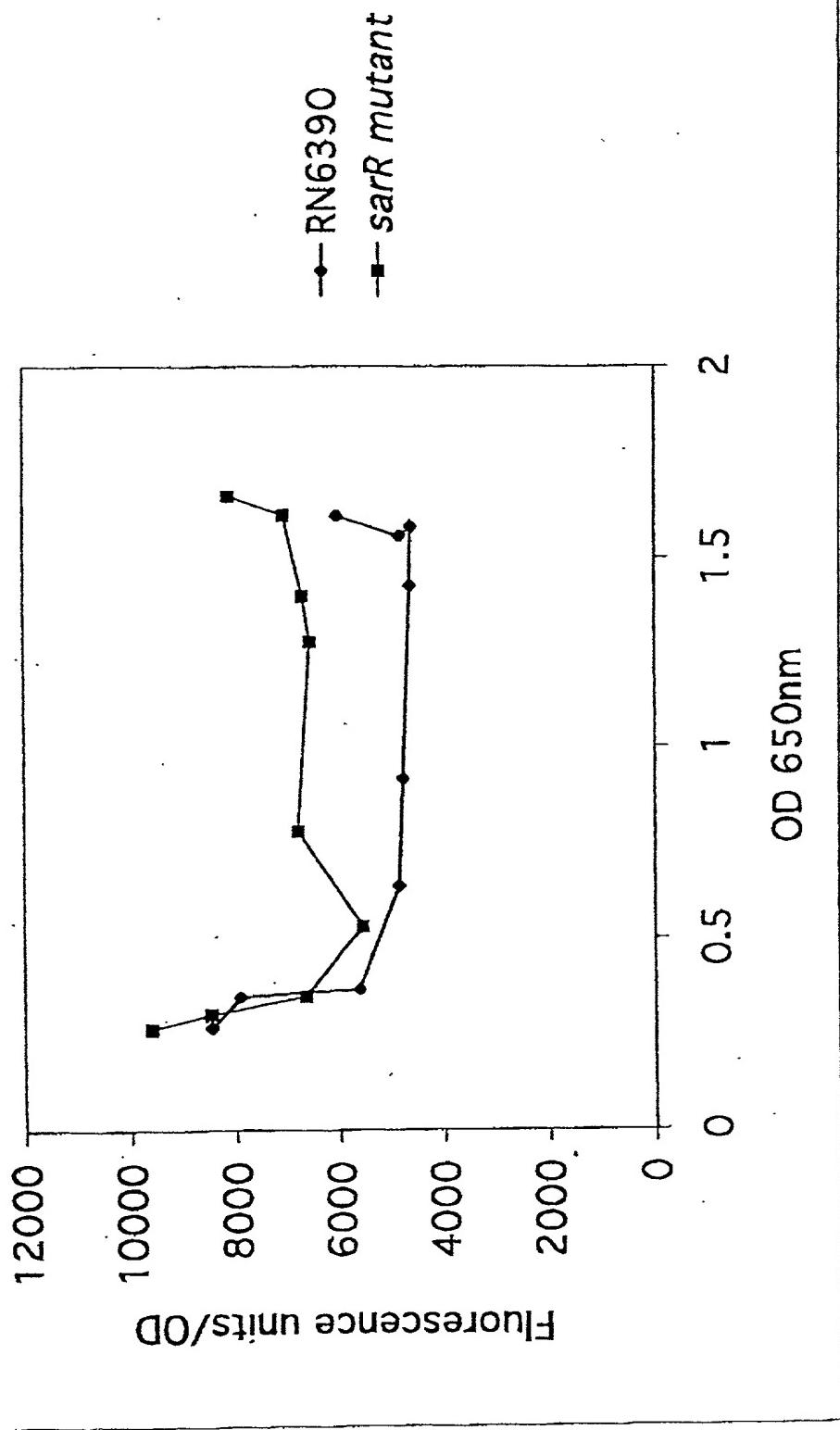


FIG. 7B

sar P2P3P1



q/15

10
15

FIG. 8A: *sar* probe

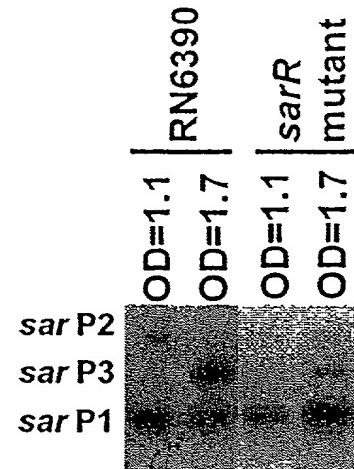


FIG. 8B: anti-SarA antibody

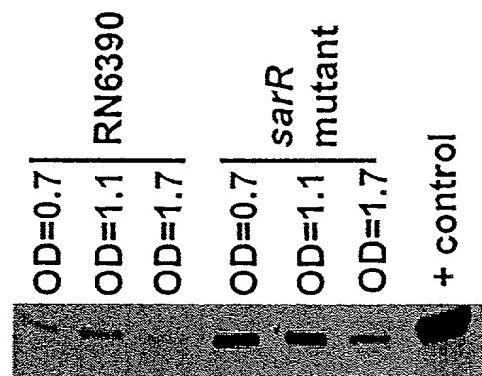
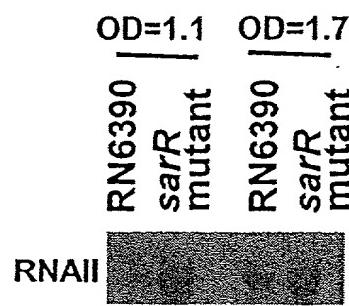


FIG. 8C: *agrA* probe



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FIG. 9A

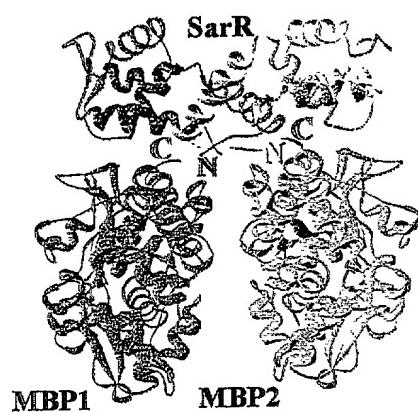
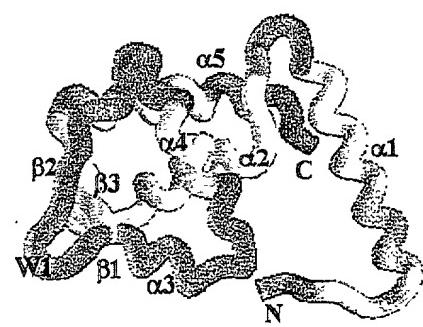
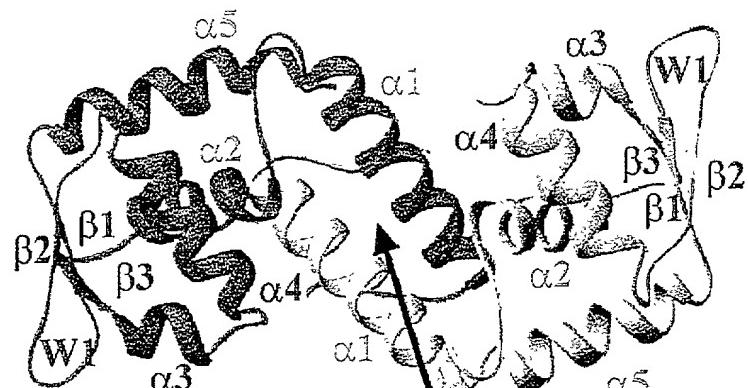


FIG. 9B



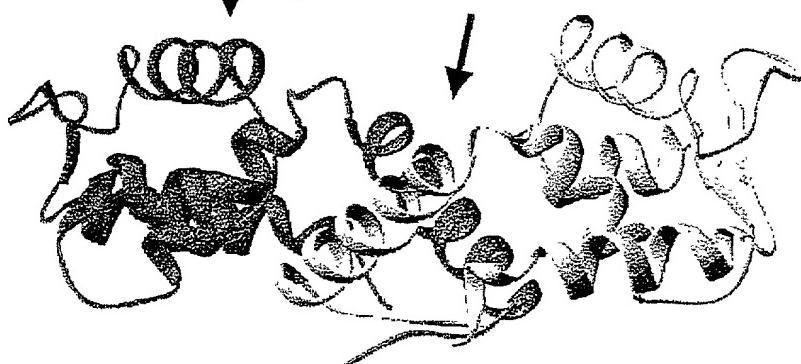
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FIG. 10A



↓ U 90° Canyon

FIG. 10B



↓ U 90°

FIG. 10C

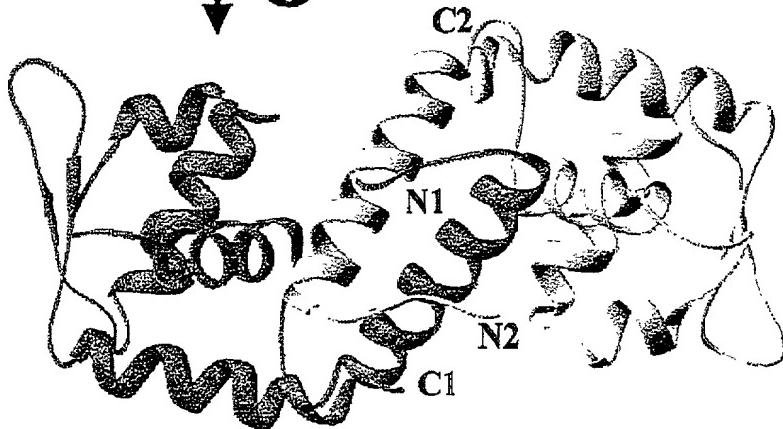


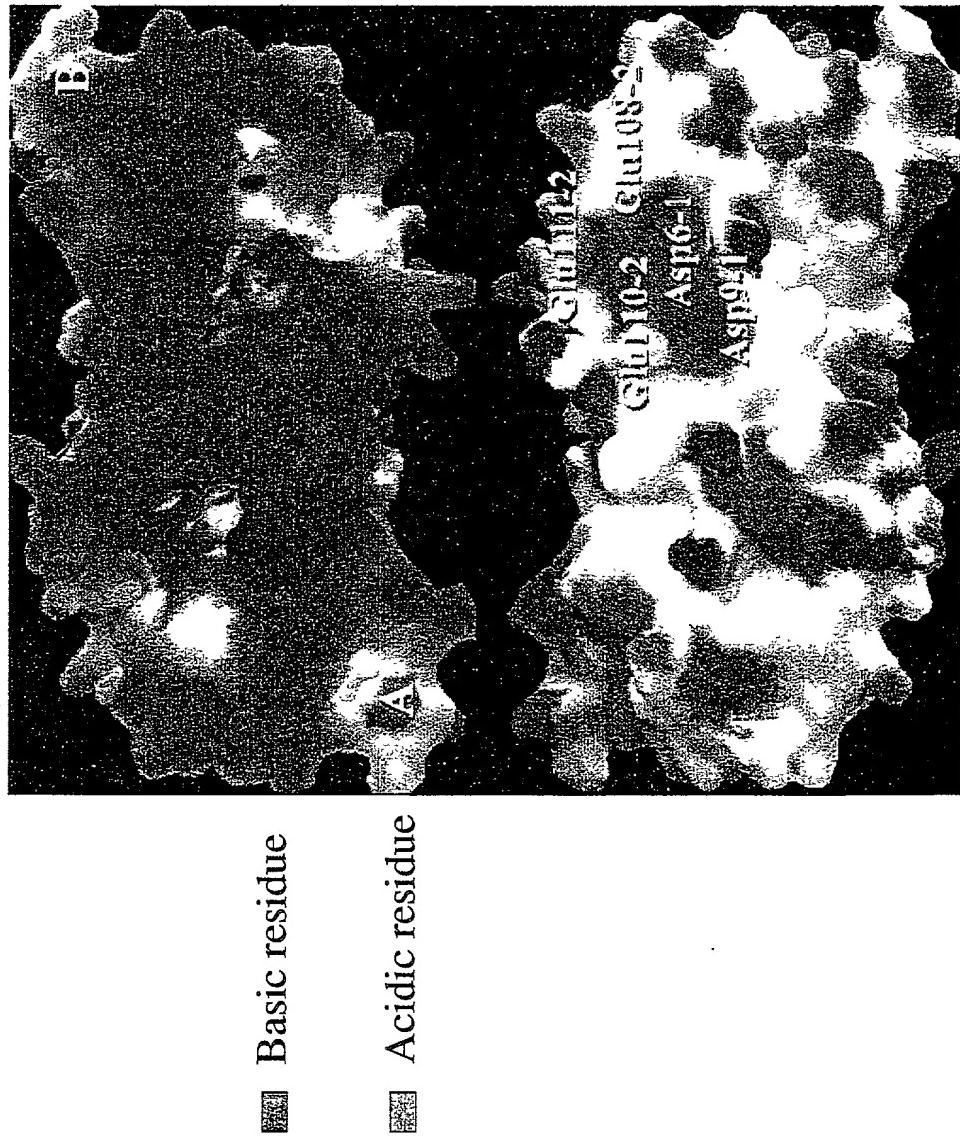
FIG. 11

Alignment of the SarA Homologs

H: residues involved in dimerization, A: acidic residues involved in activation,
 D: basic residues involved in DNA binding

FIG. 12 A & B

Electrostatic potential surface of the dimeric SarR structure



The distance from A to B is 71 Angstrom (20 nt) but the inner distance is 92 Angstrom (27 nt)

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FIG. 13A

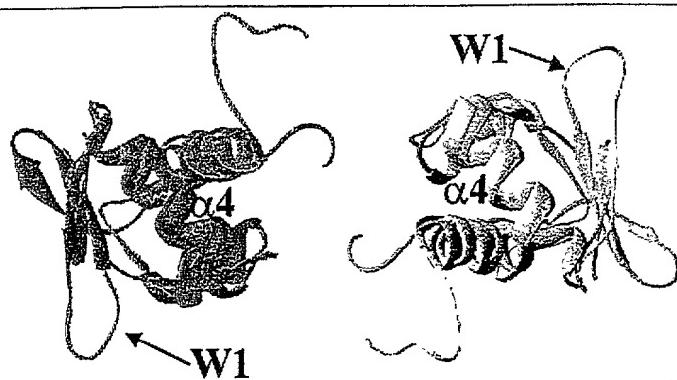


FIG. 13B

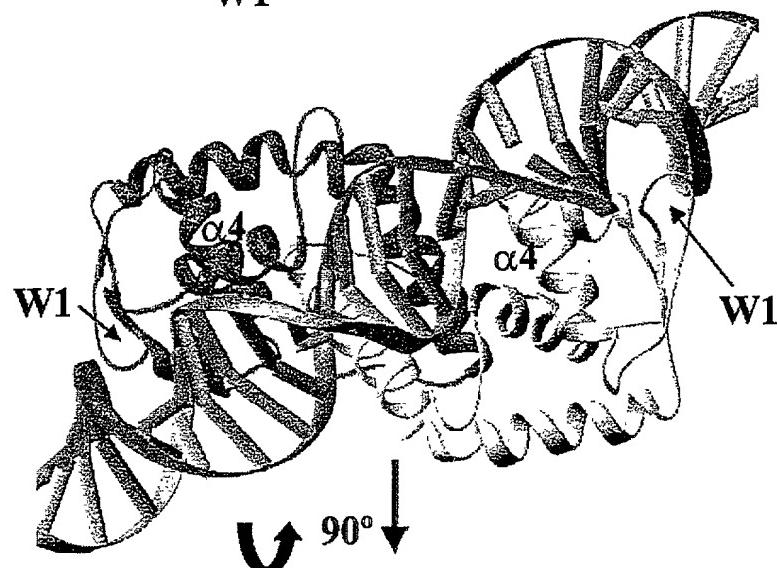


FIG. 13C

